

velocity_profiles

March 11, 2021

get started by plotting velocity profiles for simulated and fanjin data

```
[1]: from tqdm import tqdm
import os
import numpy as np
from tabulate import tabulate
import matplotlib.pyplot as plt
import matplotlib as mpl
import _fj
import plotutils
import command
import twutils
import twanalyse
import stats
import readtrack
import scipy
```

```
[2]: # paths
notename = "velocity"
pdir = notename+'/'
notedir = os.getcwd()
rsimdir = "exampledata/two_parameters/pilivar_0013.00000_k_spawn_00.50000/"
# rsimdir = "exampledata/two_parameters/pilivar_0013.00000_k_spawn_05.00000/"
simdir = os.path.join(notedir, rsimdir)
```

```
[3]: # fanjin
debug = None
debug = 100
idx, fltrs = _fj.slicehelper.load_linearized_trs('default_crawling_list', debug)
```

100%| | 100/100 [00:00<00:00, 1599.06it/s]

```
[4]: # plotting fj data
allvel = np.concatenate([twanalyse._inst_vel(tr) for tr in fltrs])
# allvel = twutils.trim_tail(allvel, 0.05)
# ax = plt.gca()
# outd = plotutils.ax_kdeplot(ax, allvel)
# fjspace, fjpde = outd['space'], outd['pde']
```

```
# ax.set_xlabel(r"velocity $\mu$ m$")
# plt.close()
```

```
[5]: # The same for simulation data
simdata = os.path.join(simdir, "data/")
trs = readtrack.trackset(ddir=simdata)

for tr in trs:
    tr._clean_bad_dt()
ltrs = [_fj.linearize(tr) for tr in trs]

simvel = np.concatenate([twanalyse._inst_vel(tr) for tr in ltrs])
# ax = plt.gca()
# # simvel = twutils.trim_tail(simvel, 0.05)
# outd = plotutils.ax_kdeplot(ax, simvel)
# simspace, simpde = outd['space'], outd['pde']
# plt.show()
```

searching for tracks with form /home/dan/usb_twitching/pili/notebook/exampledata/two_parameters/pilivar_0013.00000_k_spawn_00.50000/data/bacterium_*.dat

Comparing instantaneous velocity of linearized tracks. Tracks linearised with respect to a threshold distance 0.12 microns. Instantaneous velocity is calculated over 0.1 second interval.

```
[6]: fjevstat = stats.stats(allvel)
simvstat = stats.stats(simvel)
print("Instantaneous velocity")
print(r"Fanjin")
twutils.print_dict(fjevstat)
print(r"Simulated example [{}]".format(rsimdir))
twutils.print_dict(simvstat)
```

Instantaneous velocity

Fanjin

```
{
    "mean": 0.013918706870453825,
    "median": 0.008140381093561537,
    "std": 0.03596748602943479,
    "std_error": 3.5102878775917124e-05
}
```

Simulated example

```
[exampledata/two_parameters/pilivar_0013.00000_k_spawn_00.50000/]
{
    "mean": 0.047287128380971336,
    "median": 0.0,
    "std": 0.21345935323800733,
    "std_error": 0.0005477946688869858
}
```

```
[7]: import rtw
import txtdata

target = "../../../run/two_parameter_model/two_parameters/"
dc = rtw.DataCube(target)
lvel_mean = dc.get_local_array( rtw._make_get("lvel.mean") )
lvel_std = dc.get_local_array( rtw._make_get("lvel.std"))
fjvmean = fjvstat['mean']
fjvstd = fjvstat['std']
rel_lvel_mean = lvel_mean - fjvmean
rel_lvel_std = lvel_std - fjvstd
print("Want to analyse a 2d parameter search dataset with parameters")
table = [ [name] + base for name, base in zip(dc.pnames, list(dc.basis)) ]
print(tabulate(table, floatfmt='.2f'))
```

WARNING: parameters.thisread() did not find ./config.txt. Continuing with defaults.

WARNING: parameters.thisread() did not find ./config.txt. Continuing with defaults.

WARNING: did not find local config.txt, default params loaded

Want to analyse a 2d parameter search dataset with parameters

```
-----
-----
pilivar  1.00  2.00  2.50  3.00  3.50  4.00  7.00  13.00  20.00  50.00  100.00
k_spawn  0.10  0.50  1.00  1.50  2.00  2.50  3.00  3.50  4.00  4.50  5.00
5.50  6.00
-----
-----
```

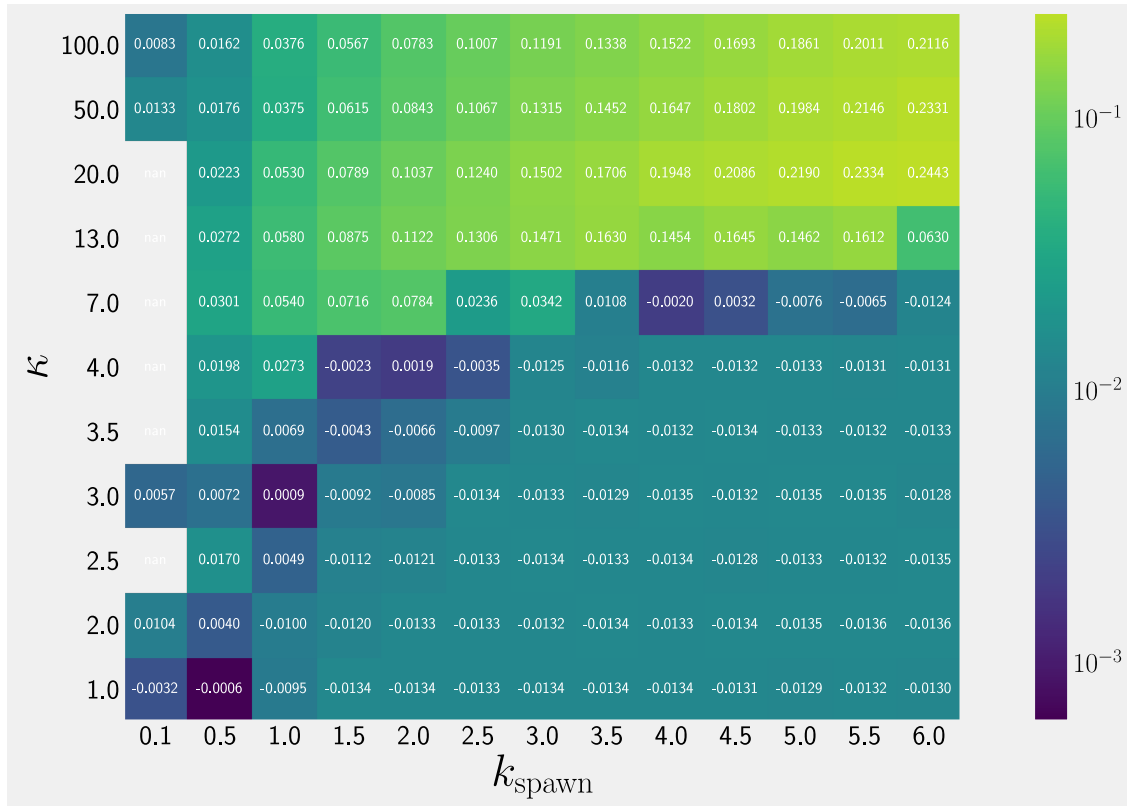
```
[8]: plt.style.use(plotutils.get_style('image'))
def relative_image(expval, localdataname):
    ax = plt.gca()
    meanget = rtw._make_get(localdataname)
    def rmeanget(ld):
        return abs(meanget(ld) - fjvmean)
    def rmeanget_an(ld):
        return meanget(ld) - fjvmean

    rtw._param_image(ax, dc, rmeanget, annotate=True,
        annotate_form=rtw.anform[localdataname], use_lognorm=True,
        ↪_getter_an=rmeanget_an)
print("mean velocity relative to FJ data")
relative_image(fjvmean, 'lvel.mean')
plt.show()

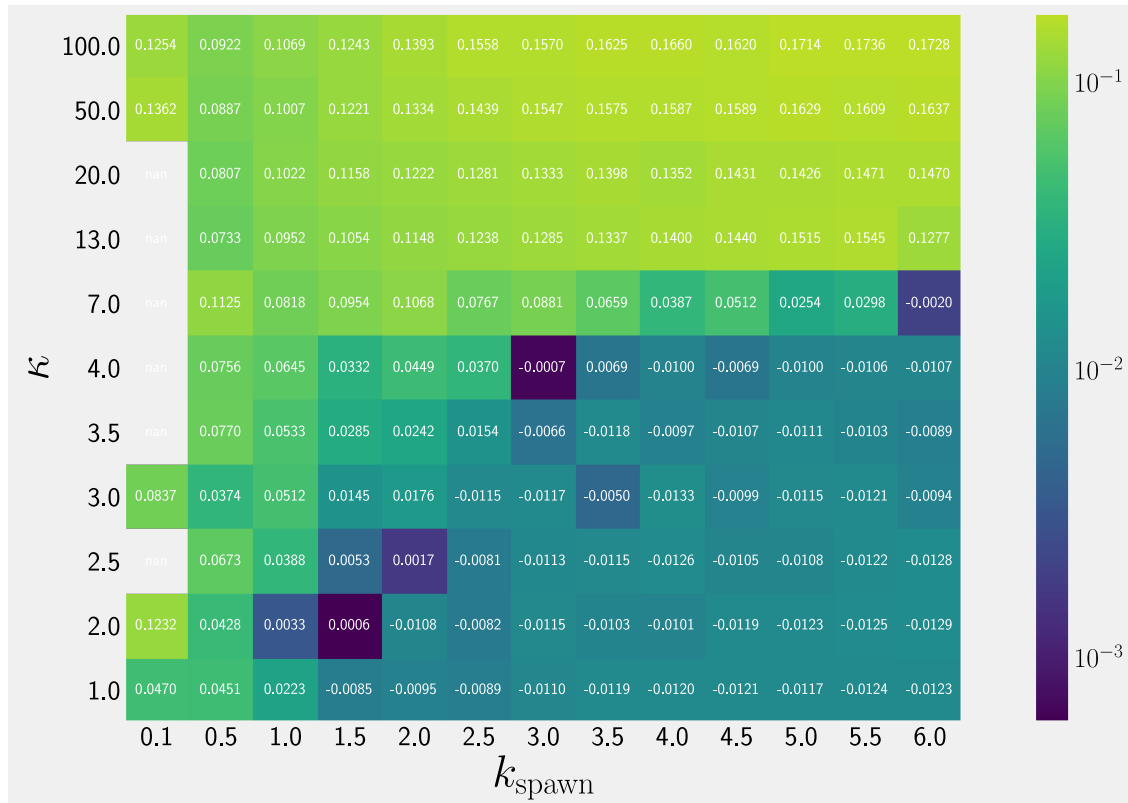
print("velocity std relative to FJ data")
relative_image(fjvstd, 'lvel.std')
```

```
plt.show()
```

mean velocity relative to FJ data



velocity std relative to FJ data



```
[9]: # so ok there are tracks with similar velocity and standard deviation but we
      ↪ have seen their shapes
      # are not so close as we might like.
      # we need to quantify this. We can use smoothing kernel to probability
      ↪ distribution and then
      # do some kind of weighted least squares but I would like a better approach.
      # go to higher order than 2nd moment? Just look at quantiles? KS statistic?
      pass
```

We want consider multiple metrics, in this case just mean velocity and standard deviation initially. Lets start by finding the pareto set.

```
[10]: # pareto
      # https://stackoverflow.com/questions/32791911/
      ↪ fast-calculation-of-pareto-front-in-python

      # Fairly fast for many datapoints, less fast for many costs, somewhat readable
      def is_pareto_efficient_simple(costs):
          """
          Find the pareto-efficient points
          :param costs: An (n_points, n_costs) array
```

```

        :return: A (n_points, ) boolean array, indicating whether each point is
        ↪ Pareto efficient
        """
        is_efficient = np.ones(costs.shape[0], dtype = bool)
        for i, c in enumerate(costs):
            if is_efficient[i]:
                # Keep any point with a lower cost
                is_efficient[is_efficient] = np.any(costs[is_efficient]<c, axis=1)
                is_efficient[i] = True # And keep self
        return is_efficient

# being above or below the experimental data is bad so take absolute value

objective_shape = rel_lvel_mean.shape
mean_objective = np.abs(rel_lvel_mean)
std_objective = np.abs(rel_lvel_std)

objectives = np.column_stack( [ mean_objective.flatten(), std_objective.
    ↪ flatten() ] )
pareto_front = is_pareto_efficient_simple(objectives)
pareto_front = pareto_front.reshape(objective_shape)
pareto_idx = np.nonzero(pareto_front)

# draw pareto front
im_front = np.full(objective_shape, 1.0)
im_front[pareto_idx] = 0.5

ax = plt.gca()

ax.set_xticks(np.arange(len(dc.slice_basis[1])))
ax.set_yticks(np.arange(len(dc.slice_basis[0])))
ax.set_xticklabels(dc.slice_basis[1])
ax.set_yticklabels(dc.slice_basis[0])
ax.set_xlabel(txtdata.prettynames.get(dc.pnames[1]))
ax.set_ylabel(txtdata.prettynames.get(dc.pnames[0]))

norm = mpl.colors.Normalize(vmin=0.0, vmax=1.0)
Image = ax.imshow(im_front, norm=norm, cmap=plt.cm.gray, origin='lower')
plt.show()

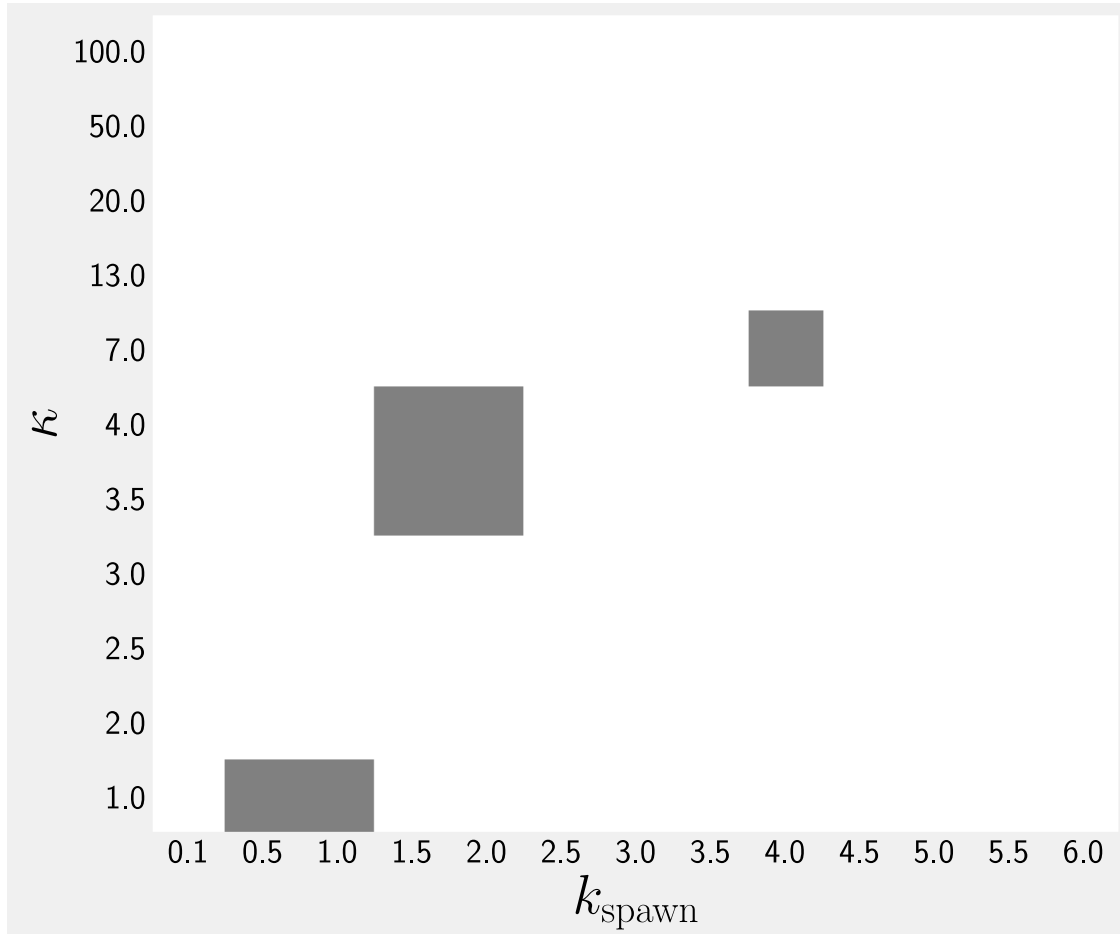
# what is the difference between min and max in pareto front
def minmax(arr):
    return np.min(arr), np.max(arr)
min_mean, max_mean = minmax(rel_lvel_mean[pareto_idx])
min_std, max_std = minmax(rel_lvel_std[pareto_idx])
amin_mean, amax_mean = minmax(mean_objective[pareto_idx])

```

```

amin_std, amax_std = minmax(std_objective[pareto_idx])
#
print('pareto front lims')
headers = ['', 'min', 'max', '', 'min', 'max']
table = tabulate([
    ['lvel mean ', min_mean, max_mean, 'absolute', amin_mean, amax_mean],
    ['lvel abs ', min_std, max_std, 'absolute', amin_std, amax_std]
], headers, floatfmt='.4f')
print(table)

```



```

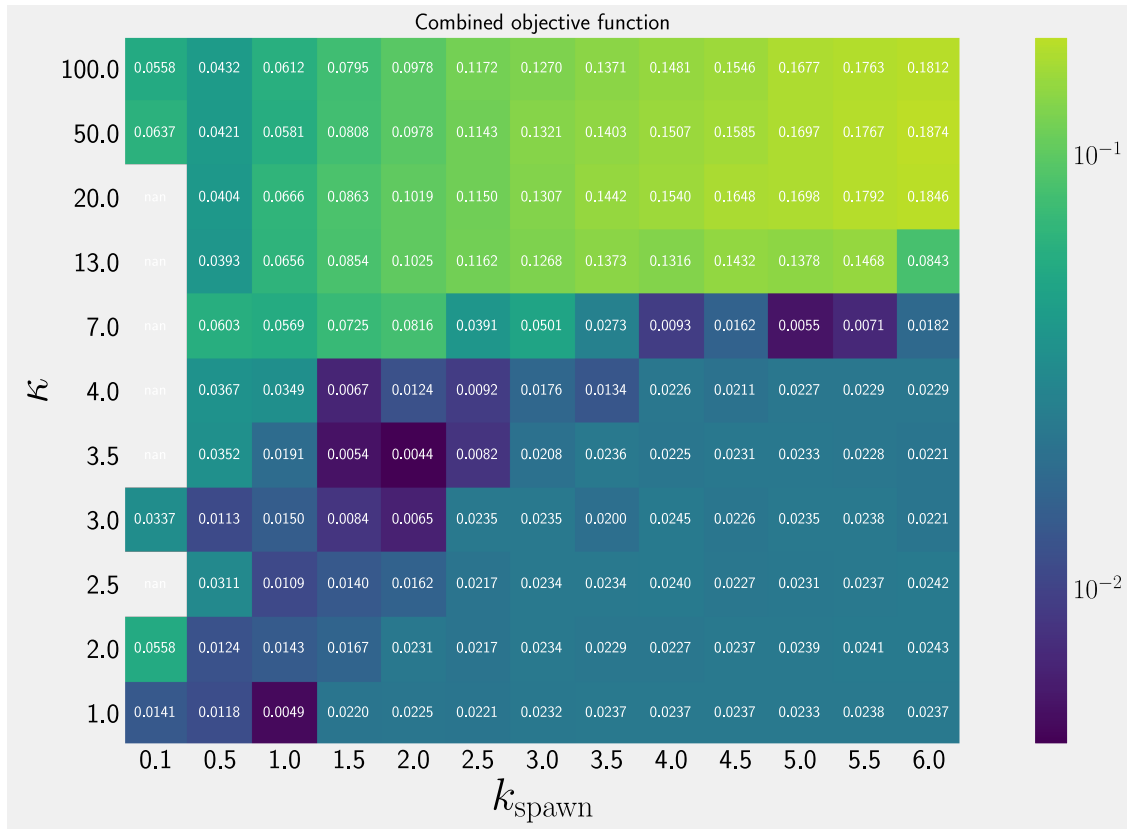
pareto front lims
-----
           min      max      absolute   min      max
lvel mean -0.0095  0.0019  absolute  0.0006  0.0095
lvel abs   0.0003  0.0230  absolute  0.0003  0.0230

```

We may want to just transform how multobjective optimisation problem into a more straightforward problem by using a linear combination of objective functions. Both parameters have similar ranges in the pareto front so in this case so lets give them equal weight.

```
[11]: # combine into a single objective function
lcojective = (mean_objective + std_objective)/2
ax = plt.gca()
rtw._data_image(ax, dc, lcojective, annotate=True, annotate_form=rtw.
    ↳anform['lvel.mean'])
ax.set_title("Combined objective function")
plt.show()

# reading off good values for both parameters
eye_values = [[1.0, 1.0], [3.5, 2.0], [7.0, 5.0]]
eye_table = tabulate(eye_values, dc.pnames, floatfmt='.2f')
print('Hand picking some of the parameters from across the parameter space.')
print(eye_table)
```



Hand picking some of the parameters from across the parameter space.

pilivar	k_spawn
1.00	1.00
3.50	2.00
7.00	5.00


```
[12]: # retrieve simulation index and path
eye_idx = [dc.find_index(xy) for xy in eye_values]
eye_dir = [dc.dircube[tuple(idx)] for idx in eye_idx]

# load linearized velocities
# for i, eye
lsimvel = []
for i, idx in enumerate(eye_idx):
    simdir = eye_dir[i]
    print('loading data from ', simdir)
    simdata = os.path.join(simdir, "data/")
    trs = readtrack.trackset(ddir=simdata)
    for tr in trs:
        tr._clean_bad_dt()
    ltrs = [_fj.linearize(tr) for tr in trs]
    track_vel = [twanalyse._inst_vel(tr) for tr in ltrs]
    simvel = np.concatenate(track_vel)
    # simvel = twutils.trim_tail(simvel, 0.05)
    lsimvel.append(simvel)
```

```
loading data from ../../run/two_parameter_model/two_parameters/pilivar_0001.000
00_k_spawn_01.00000/
searching for tracks with form ../../run/two_parameter_model/two_parameters/pili
var_0001.00000_k_spawn_01.00000/data/bacterium*.dat
loading data from ../../run/two_parameter_model/two_parameters/pilivar_0003.500
00_k_spawn_02.00000/
searching for tracks with form ../../run/two_parameter_model/two_parameters/pili
var_0003.50000_k_spawn_02.00000/data/bacterium*.dat
loading data from ../../run/two_parameter_model/two_parameters/pilivar_0007.000
00_k_spawn_05.00000/
searching for tracks with form ../../run/two_parameter_model/two_parameters/pili
var_0007.00000_k_spawn_05.00000/data/bacterium*.dat
```

```
[13]: # check the bounds
table = []
for i, idx in enumerate(eye_idx):
    simvel = lsimvel[i]
    row = [os.path.basename(os.path.normpath(eye_dir[i]))]
    row.extend(np.quantile(simvel, [0.0, 0.25, 0.5, 0.75, 1.0]).tolist())
    table.append(row)
table.append(['Fanjin', *np.quantile(allvel, [0.0, 0.25, 0.5, 0.75, 1.0]).
    ↳tolist()])
print(tabulate(table, headers=['simulation path', 'min', '1st', '2nd', '3rd', '
    ↳max'], floatfmt='.4f'))
```

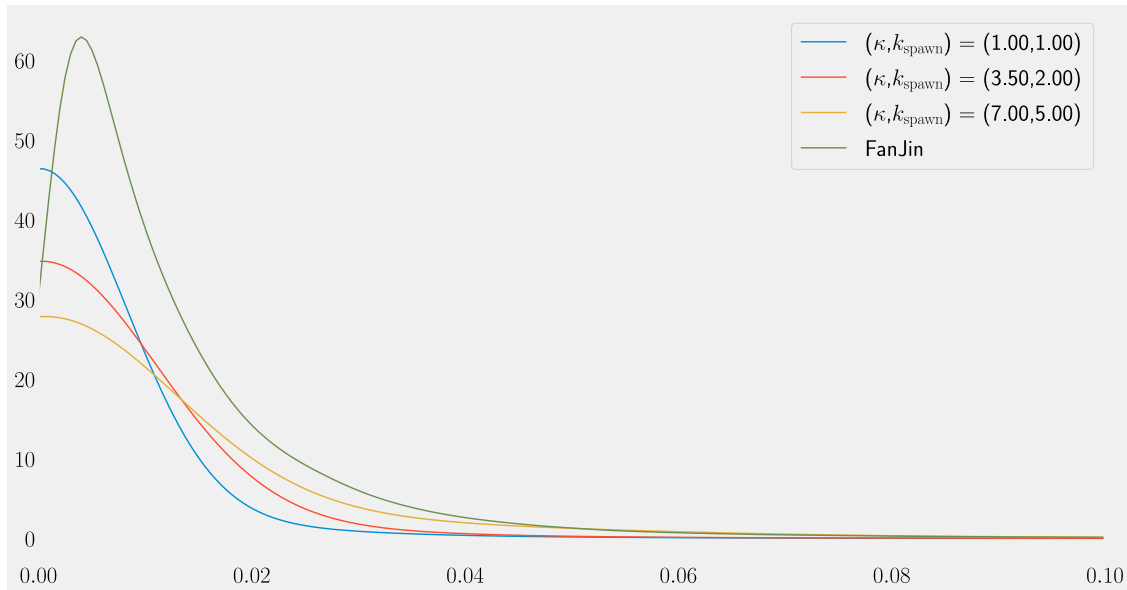
simulation path	min	1st	2nd	3rd	max
-----------------	-----	-----	-----	-----	-----

pilivar_0001.00000_k_spawn_01.00000	0.0000	0.0000	0.0000	0.0000	3.7041
pilivar_0003.50000_k_spawn_02.00000	0.0000	0.0000	0.0000	0.0000	3.8298
pilivar_0007.00000_k_spawn_05.00000	0.0000	0.0000	0.0000	0.0000	4.0678
Fanjin	0.0000	0.0038	0.0081	0.0160	10.1829

```
[14]: # We notice immediately from printing the 1st, 2nd and 3rd quantiles that these
      ↪ distributions
      # are not all that similar. Simulated tracks in this dataset spend a large
      ↪ portion of the time
      # idling at close to 0 velocity. This is even after linearising the trajectory.
      ↪ (Worth checking again)
```

```
[15]: ax = plt.gca()
      plt.style.use(plotutils.get_style('jupyter'))
      handle = []
      label = []
      xlims = (0, 0.1)
      use_hist = False
      for i, simvel in enumerate(lsimvel):
          outd = plotutils.ax_kdeplot(ax, simvel, xlims=xlims, hist=use_hist)
          handle.append(outd['handle'])
          label.append("({}, {}) = ({:4.2f}, {:4.2f})".format(*dc.prettynames(),
          ↪ *eye_values[i]))
      outd = plotutils.ax_kdeplot(ax, allvel, xlims=xlims, hist=use_hist)
      label.append('FanJin')
      handle.append(outd['handle'])
      ax.legend(handle, label)
      ax.set_xlim(xlims)
      plt.show()
```

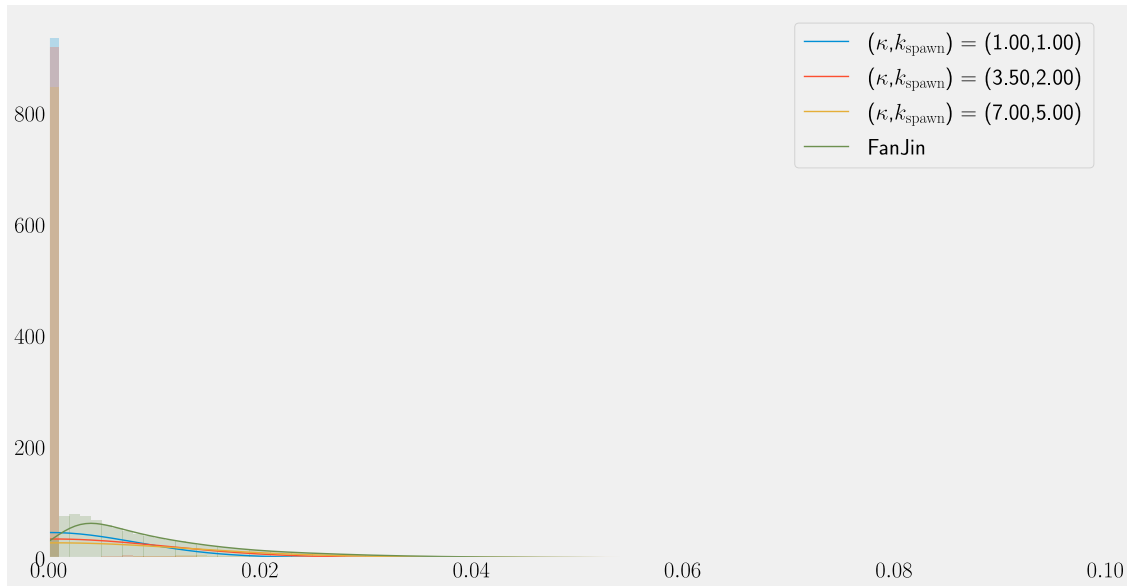
Computing pdf with 195215 data points at resolution 200
 Computing pdf with 199542 data points at resolution 200
 Computing pdf with 199961 data points at resolution 200
 Computing pdf with 1049869 data points at resolution 200



These are probability distributions and should all have the same area. Most likely the kernel we use to compute probability density is not reflected at $x = 0$ but this library doesn't give the option to change that. It's always important to plot a straightforward histogram.

```
[16]: ax = plt.gca()
plt.style.use(plotutils.get_style('jupyter'))
handle = []
label = []
xlims = (0, 0.1)
use_hist = True
for i, simvel in enumerate(lsimvel):
    outd = plotutils.ax_kdeplot(ax, simvel, xlims=xlims, hist=use_hist)
    handle.append(outd['handle'])
    label.append("({}, {}) = ({}:4.2f, {}:4.2f)".format(*dc.prettynames(),
↪ *eye_values[i]))
outd = plotutils.ax_kdeplot(ax, allvel, xlims=xlims, hist=use_hist)
label.append('FanJin')
handle.append(outd['handle'])
ax.legend(handle, label)
ax.set_xlim(xlims)
plt.show()
```

```
Computing pdf with 195215 data points at resolution 200
Computing pdf with 199542 data points at resolution 200
Computing pdf with 199961 data points at resolution 200
Computing pdf with 1049869 data points at resolution 200
```



A histogram shows the problem with simulated tracks spending large amounts of time stationary. It's been a while so I do need to test the code again for bugs and before I analyse the trajectories to understand why this happens. it's also worth noting that standard deviation of velocity velocity is not very useful here because the simulated distribution is so skewed.